Resource Summary Report

Generated by dkNET on May 19, 2025

BSA4Yeast

RRID:SCR_017113

Type: Tool

Proper Citation

BSA4Yeast (RRID:SCR_017113)

Resource Information

URL: https://bsa4yeast.lcsb.uni.lu

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Description: Web application for Quantitative Trait Loci mapping via bulk segregant analysis of yeast sequencing data. Application provides automated data processing, annotations, and web interface to explore identified QTLs.

Synonyms: Bulk Segregant Analysis for Yeast

Resource Type: data access protocol, service resource, production service resource, data analysis service, analysis service resource, web service, software resource

Keywords: quantitative, trait, loci, mapping, bulk, segregant, analysis, yeast, sequencing, data, processing, annotation, bio.tools

Funding: Fonds Nationale de la Recherche (FNR) Luxembourg

Availability: Free, Freely available, Registration required

Resource Name: BSA4Yeast

Resource ID: SCR_017113

Alternate IDs: biotools:BSA4Yeast

Alternate URLs: https://bio.tools/BSA4Yeast

Record Creation Time: 20220129T080333+0000

Record Last Update: 20250517T060312+0000

Ratings and Alerts

No rating or validation information has been found for BSA4Yeast.

No alerts have been found for BSA4Yeast.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Zhang Z, et al. (2019) BSA4Yeast: Web-based quantitative trait locus linkage analysis and bulk segregant analysis of yeast sequencing data. GigaScience, 8(6).